



PCT10

**Does Not Comply
Corrected Diskette Needed**

RAW SEQUENCE LISTING

PATENT APPLICATION: US/10/019,065

DATE: 07/16/2002

TIME: 13:52:22

Input Set : A:\EP.txt

Output Set: N:\CRF3\07162002\J019065.raw

PatentIn 2.0 "bug". See
p. 7 for explanation

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804 <213> ORGANISM: human
806 <400> SEQUENCE: 33

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825	Arg	Asn	Ala	Ala	Gly	Ser	Ala	His	Arg	His	Val	Thr	Leu	His	Val	His
826				100					105					110		
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829				115				120					125			
831	Asn	Asn	Pro	Ile	Leu	Leu	Pro	Cys	Glu	Ala	Thr	Gly	Thr	Pro	Ser	Pro
832		130					135					140				
834	Phe	Ile	Thr	Trp	Gln	Lys	Glu	Gly	Ile	Asn	Val	Asn	Thr	Ser	Gly	Arg
835	145				150					155					160	
837	Asn	His	Ala	Val	Leu	Pro	Ser	Gly	Gly	Leu	Gln	Ile	Ser	Arg	Ala	Val
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841				180					185					190		
843	Thr	Ala	Leu	Gly	Lys	Ile	Lys	Leu	Asn	Val	Gln	Val	Pro	Pro	Val	Ile
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849	Leu	Ser	Cys	Glu	Ala	Asp	Gly	Leu	Pro	Pro	Pro	Asp	Ile	Thr	Trp	His
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852	Lys	Asp	Gly	Arg	Ala	Ile	Val	Glu	Ser	Ile	Arg	Gln	Arg	Val	Leu	Ser

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856 Ser Gly Ser Leu Gln Ile Ala Phe Val Gln Pro Gly Asp Ala Gly His
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859 Tyr Thr Cys Met Ala Ala Asn Val Ala Gly Ser Ser Ser Thr Ser Thr
860          275          280          285
862 Lys Leu Thr Val His Val Pro Pro Arg Ile Arg Ser Thr Lys Gly His
863          290          295          300
865 Tyr Thr Val Asn Glu Asn Ser Gln Ala Ile Leu Pro Cys Val Ala Asp
866 305          310          315          320
868 Gly Ile Pro Thr Pro Ala Ile Asn Trp Lys Lys Asp Asn Val Leu Leu
869          325          330          335
871 Ala Asn Leu Leu Gly Lys Tyr Thr Ala Glu Pro Tyr Gly Glu Leu Ile
872          340          345          350
874 Leu Glu Asn Val Val Leu Glu Asp Ser Gly Phe Tyr Thr Cys Val Ala
875          355          360          365
877 Asn Asn Ala Ala Gly Glu Asp Thr His Thr Val Ser Leu Thr Val His
878          370          375          380
880 Val Leu Pro Thr Phe Thr Glu Leu Pro Gly Asp Val Ser Leu Asn Lys
881 385          390          395          400
883 Gly Glu Gln Leu Arg Leu Ser Cys Lys Ala Thr Gly Ile Pro Leu Pro
884          405          410          415
886 Lys Leu Thr Trp Thr Phe Asn Asn Asn Ile Ile Pro Ala His Phe Asp
887          420          425          430
889 Ser Val Asn Gly His Ser Glu Leu Val Ile Glu Arg Val Ser Lys Glu
890          435          440          445
892 Asp Ser Gly Thr Tyr Val Cys Thr Ala Glu Asn Ser Val Gly Phe Val
893          450          455          460
895 Lys Ala Ile Gly Phe Val Tyr Val Lys Glu Pro Pro Val Phe Lys Gly
896 465          470          475          480
898 Asp Tyr Pro Ser Asn Trp Ile Glu Pro Leu Gly Gly Asn Ala Ile Leu
899          485          490          495
901 Asn Cys Glu Val Lys Gly Asp Pro Thr Pro Thr Ile Gln Trp Asn Arg
902          500          505          510
904 Lys Gly Val Asp Ile Glu Ile Ser His Arg Ile Arg Gln Leu Gly Asn
905          515          520          525
907 Gly Ser Leu Ala Ile Tyr Gly Thr Val Asn Glu Asp Ala Gly Asp Tyr
908          530          535          540
910 Thr Cys Val Ala Thr Asn Glu Ala Gly Val Val Glu Arg Ser Met Ser
911 545          550          555          560
913 Leu Thr Leu Arg Ser Pro Pro Ile Ile Thr Leu Glu Pro Val Glu Thr
914          565          570          575
916 Val Ile Asn Ala Gly Gly Lys Ile Ile Leu Asn Cys Gln Ala Thr Gly
917          580          585          590
919 Glu Pro Gln Pro Thr Ile Thr Trp Ser Arg Gln Gly His Ser Ile Ser
920          595          600          605
922 Trp Asp Asp Arg Val Asn Val Leu Ser Asn Asn Ser Leu Tyr Ile Ala
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925 Asp Ala Gln Lys Glu Asp Thr Ser Glu Phe Glu Cys Val Ala Arg Asn
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934 Gly Lys Gly Ile Gln Lys Arg Ser Arg Leu Cys Asn Gln Pro Leu Pro
935           675           680           685
937 Ala Asn Gly Gly Lys Pro Cys Gln Gly Ser Asp Leu Glu Met Arg Asn
938           690           695           700
940 Cys Gln Asn Lys Pro Cys Pro Val Asp Gly Ser Trp Ser Glu Trp Ser
941 705           710           715           720
943 Leu Trp Glu Glu Cys Thr Arg Ser Cys Gly Arg Gly Asn Gln Thr Arg
944           725           730           735
946 Thr Arg Thr Cys Asn Asn Pro Ser Val Gln His Gly Gly Arg Pro Cys
947           740           745           750
949 Glu Gly Asn Ala Val Glu Ile Ile Met Cys Asn Ile Arg Pro Cys Pro
950           755           760           765
952 Val His Gly Ala Trp Ser Ala Trp Gln Pro Trp Gly Thr Cys Ser Glu
953           770           775           780
955 Ser Cys Gly Lys Gly Thr Gln Thr Arg Ala Arg Leu Cys Asn Asn Pro
956 785           790           795           800
958 Pro Pro Ala Phe Gly Gly Ser Tyr Cys Asp Gly Ala Glu Thr Gln Met
959           805           810           815
961 Gln Val Cys Asn Glu Arg Asn Cys Pro Ile His Gly Lys Trp Ala Thr
962           820           825           830
964 Trp Ala Ser Trp Ser Ala Cys Ser Val Ser Cys Gly Gly Gly Ala Arg
965           835           840           845
967 Gln Arg Thr Arg Gly Cys Ser Asp Pro Val Pro Gln Tyr Gly Gly Arg
968           850           855           860
970 Lys Cys Glu Gly Ser Asp Val Gln Ser Asp Phe Cys Asn Ser Asp Pro
971 865           870           875           880
973 Cys Pro Thr His Gly Asn Trp Ser Pro Trp Ser Gly Trp Gly Thr Cys
974           885           890           895
976 Ser Arg Thr Cys Asn Gly Gly Gln Met Arg Arg Tyr Arg Thr Cys Asp
977           900           905           910
979 Asn Pro Pro Pro Ser Asn Gly Gly Arg Ala Cys Gly Gly Pro Asp Ser
980           915           920           925
982 Gln Ile Gln Arg Cys Asn Thr Asp Met Cys Pro Val Asp Gly Ser Trp
983           930           935           940
985 Gly Ser Trp His Ser Trp Ser Gln Cys Ser Ala Ser Cys Gly Gly Gly
986 945           950           955           960
988 Glu Lys Thr Arg Lys Arg Leu Cys Asp His Pro Val Pro Val Lys Gly
989           965           970           975
991 Gly Arg Pro Cys Pro Gly Asp Thr Thr Gln Val Thr Arg Cys Asn Val
992           980           985           990
994 Gln Ala Cys Pro Gly Gly Pro Gln Arg Ala Arg Gly Ser Val Ile Gly
995           995           1000           1005
997 Asn Ile Asn Asp Val Glu Phe Gly Ile Ala Phe Leu Asn Ala Thr Ile
998 1010           1015           1020
1000 Thr Asp Ser Pro Asn Ser Asp Thr Arg Ile Ile Arg Ala Lys Ile Thr

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1007          1060          1065          1070
1009 Asn Gly Phe Thr Leu Thr Asn Ala Val Phe Lys Arg Glu Thr Gln Val
1010          1075          1080          1085
1012 Glu Phe Ala Thr Gly Glu Ile Leu Gln Met Ser His Ile Ala Arg Gly
1013          1090          1095          1100
1015 Leu Asp Ser Asp Gly Ser Leu Leu Leu Asp Ile Val Val Ser Gly Tyr
1016 1105          1110          1115          1120
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1019          1125          1130          1135
1021 Glu Asp Tyr Ile Gln Thr Gly Pro Gly Gln Leu Tyr Ala Tyr Ser Thr
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1024 Arg Leu Phe Thr Ile Asp Gly Ile Ser Ile Pro Tyr Thr Trp Asn His
1025          1155          1160          1165
1027 Thr Val Phe Tyr Asp Gln Ala Gln Gly Arg Met Pro Phe Leu Val Glu
1028          1170          1175          1180
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1037          1220          1225          1230
1039 Ala Asp Glu Asp Glu Cys Ala Ala Gly Asn Pro Cys Ser His Ser Cys
1040          1235          1240          1245
1042 His Asn Ala Met Gly Thr Tyr Tyr Cys Ser Cys Pro Lys Gly Leu Thr
1043          1250          1255          1260
1045 Ile Ala Ala Asp Gly Arg Thr Cys Gln Asp Ile Asp Glu Cys Ala Leu
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1048 Gly Arg His Thr Cys His Ala Gly Gln Asp Cys Asp Asn Thr Ile Gly
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PATENT APPLICATION: US/10/019,065

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PatentIn 2.0 "bug":

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VERIFICATION SUMMARY

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PATENT APPLICATION: US/10/019,065

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M:254 Repeated in SeqNo=34
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